Assignment 5: Data Visualization

Summer Heschong

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OVERVIEW

This exercise accompanies the lessons in Environmental Data Analytics on Data Visualization

Directions

- 1. Rename this file <FirstLast>_A05_DataVisualization.Rmd (replacing <FirstLast> with your first and last name).
- 2. Change "Student Name" on line 3 (above) with your name.
- 3. Work through the steps, **creating code and output** that fulfill each instruction.
- 4. Be sure your code is tidy; use line breaks to ensure your code fits in the knitted output.
- 5. Be sure to **answer the questions** in this assignment document.
- 6. When you have completed the assignment, **Knit** the text and code into a single PDF file.

Set up your session

- 1. Set up your session. Load the tidyverse, lubridate, here & cowplot packages, and verify your home directory. Read in the NTL-LTER processed data files for nutrients and chemistry/physics for Peter and Paul Lakes (use the tidy NTL-LTER_Lake_Chemistry_Nutrients_PeterPaul_Processed.csv version in the Processed_KEY folder) and the processed data file for the Niwot Ridge litter dataset (use the NEON_NIWO_Litter_mass_trap_Processed.csv version, again from the Processed_KEY folder).
- 2. Make sure R is reading dates as date format; if not change the format to date.

```
library(tidyverse); library(lubridate); library(here); library(cowplot)
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr
              1.1.4
                        v readr
                                    2.1.5
## v forcats
              1.0.0
                                    1.5.1
                        v stringr
## v ggplot2
              3.5.1
                        v tibble
                                    3.2.1
## v lubridate 1.9.3
                        v tidyr
                                    1.3.1
## v purrr
              1.0.2
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                    masks stats::lag()
## i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become error
```

```
## here() starts at /home/guest/EDE_Fall2024
##
##
## Attaching package: 'cowplot'
##
##
## The following object is masked from 'package:lubridate':
##
##
       stamp
getwd()
## [1] "/home/guest/EDE_Fall2024"
here()
## [1] "/home/guest/EDE_Fall2024"
PeterPaul.chem.nutrients <- read.csv(here(</pre>
  'Data/Processed_KEY/NTL-LTER_Lake_Chemistry_Nutrients_PeterPaul_Processed.csv'),
  stringsAsFactors = TRUE)
Litter <- read.csv(here(</pre>
  'Data/Processed_KEY/NEON_NIWO_Litter_mass_trap_Processed.csv'),
  stringsAsFactors = TRUE)
#2
PeterPaul.chem.nutrients\sampledate <- ymd(PeterPaul.chem.nutrients\sampledate)
Litter$collectDate <- ymd(Litter$collectDate)</pre>
```

Define your theme

- 3. Build a theme and set it as your default theme. Customize the look of at least two of the following:
- Plot background
- Plot title
- Axis labels
- Axis ticks/gridlines
- Legend

```
#3
my_theme <-
theme(
   panel.background = element_rect(fill = 'lightblue'),
   plot.title = element_text(
      hjust = 0.5,
      size = 16,
      face= 'bold',
      colour = 'black')
)</pre>
```

Create graphs

For numbers 4-7, create ggplot graphs and adjust aesthetics to follow best practices for data visualization. Ensure your theme, color palettes, axes, and additional aesthetics are edited accordingly.

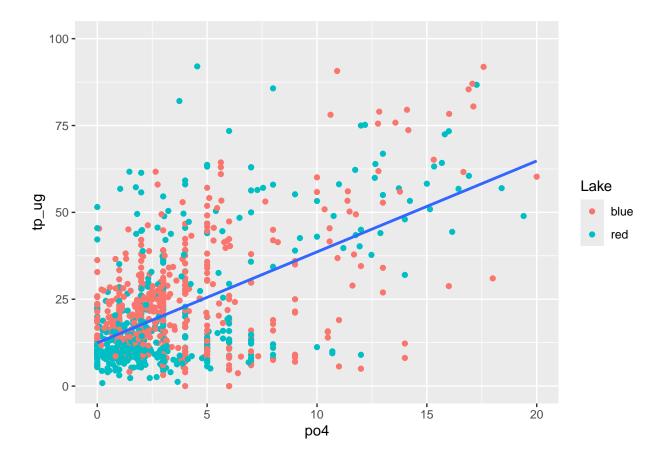
4. [NTL-LTER] Plot total phosphorus (tp_ug) by phosphate (po4), with separate aesthetics for Peter and Paul lakes. Add line(s) of best fit using the lm method. Adjust your axes to hide extreme values (hint: change the limits using xlim() and/or ylim()).

```
#4
tpbypo4_plot <- PeterPaul.chem.nutrients %>%
    ggplot(aes(x=po4, y=tp_ug))+
    geom_point(aes(color=ifelse(lakename == 'Peter Lake', 'blue', 'red'))) +
    geom_smooth(method = 'lm',se=FALSE) +
    xlim(0, 20) +
    ylim(0, 100) +
    labs(color='Lake') #I can't figure out how to change the labels
#of the blue and red dots in the legend
tpbypo4_plot

## 'geom_smooth()' using formula = 'y ~ x'

## Warning: Removed 21986 rows containing non-finite outside the scale range
## ('stat_smooth()').

## Warning: Removed 21986 rows containing missing values or values outside the scale range
## ('geom_point()').
```



5. [NTL-LTER] Make three separate boxplots of (a) temperature, (b) TP, and (c) TN, with month as the x axis and lake as a color aesthetic. Then, create a cowplot that combines the three graphs. Make sure that only one legend is present and that graph axes are aligned.

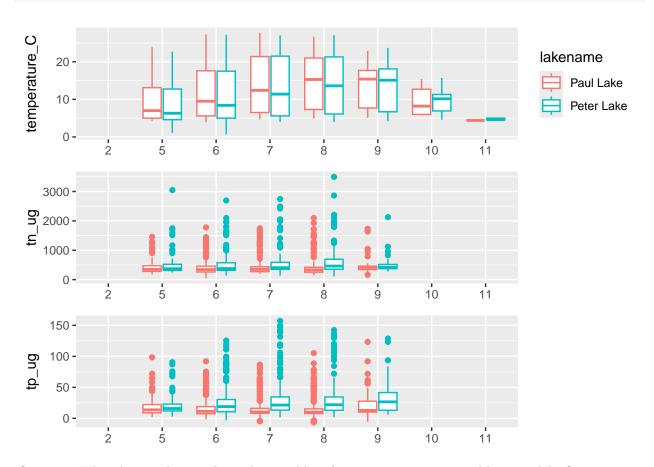
Tips: * Recall the discussion on factors in the lab section as it may be helpful here. * Setting an axis title in your theme to element_blank() removes the axis title (useful when multiple, aligned plots use the same axis values) * Setting a legend's position to "none" will remove the legend from a plot. * Individual plots can have different sizes when combined using cowplot.

```
## Warning: Removed 3566 rows containing non-finite outside the scale range
## ('stat_boxplot()').
```

```
## Warning: Removed 21583 rows containing non-finite outside the scale range
## ('stat_boxplot()').
```

Warning: Removed 20729 rows containing non-finite outside the scale range
('stat_boxplot()').

TpTnTemp_boxplot

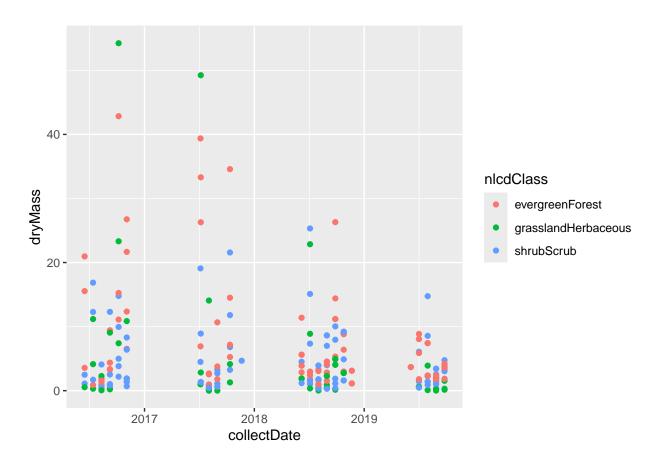


Question: What do you observe about the variables of interest over seasons and between lakes?

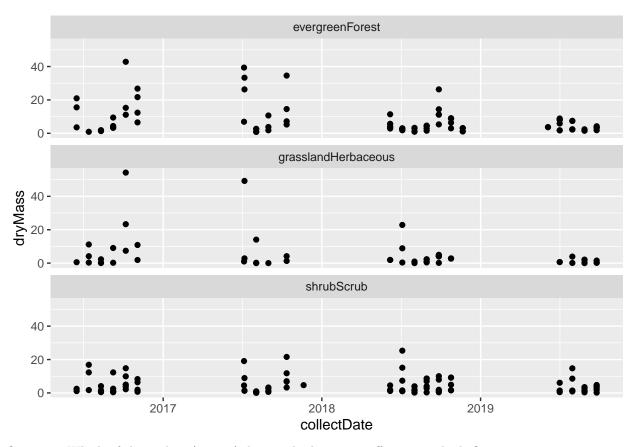
Answer: The temperature is highest in late summer for both of the lakes. To and Tp are slightly higher in Peter Lake than Paul Lake.

- 6. [Niwot Ridge] Plot a subset of the litter dataset by displaying only the "Needles" functional group. Plot the dry mass of needle litter by date and separate by NLCD class with a color aesthetic. (no need to adjust the name of each land use)
- 7. [Niwot Ridge] Now, plot the same plot but with NLCD classes separated into three facets rather than separated by color.

```
#6
litter_plotcolor <- Litter %>%
  filter(functionalGroup=='Needles') %>%
  ggplot(aes(x=collectDate, y=dryMass, color=nlcdClass)) +
  geom_point()
litter_plotcolor
```



```
#7
litter_plotfacet <- Litter %>%
  filter(functionalGroup=='Needles') %>%
  ggplot(aes(x=collectDate, y=dryMass)) +
  geom_point() +
  facet_wrap(vars(nlcdClass), nrow=3)
litter_plotfacet
```



Question: Which of these plots (6 vs. 7) do you think is more effective, and why?

Answer:I think that plot 7 is more effective because in plot 6 the points overlap and hide each other so it's hard to see all the data.